



SEQ **CLONE**

SEQ	COLORS	LC-COR1	LC-COR2	LC-COR3	RC-COR1	RC-COR2	RC-COR3	CUT	S	PL	Ramos	TF-1	2d	Marker
1	A2c	TLSTOVSSTVLA	HSEELHQS	ADARAGSSTVW	NYMT	ILSSGDB--DLSNAX	DDEGIDYDNGITLL	+	+	+	-	-		S
2	G12.1c	QASSTVS---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	YTPVYGEI--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-	CD13	L
17	B4.2a	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	TYGV	NDPILYWD--YDLYWD	DALYDASBQ---YDRL	+	+	+	-	-		L
3	E1c	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-	CD23	S
18	F2d	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	TYGV	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
4	E5e	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
5	H6.2b	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
19	G10.1	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
6	D11.1c	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-	CD23	S
20	A5.2c	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
7	F1d	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
8	F1e	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
21	E4.2	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
9	E2c	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
34	A9c	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
10	E11e	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		S
11	A1.1	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
12	F3.2	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
22	F10b	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
23	F7a	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
13	F5b	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
24	G12b	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
14	D2.1b	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
25	D1.1	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
15	D2.2a	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L
16	D2.2b	QASSTIN---YLA	QASSTL---ES	QSGOTYBA---GLT	STYLS	ILSSGDB--YDLYWD	DVLYSVYD---YDRL	+	+	+	-	-		L

CLONE: designation of representative clone for sequence

HC: Ig heavy chain

Expression pattern: binding of scv antibodies to primary human cere-

RL: non-Hodgkin's lymphoma cell line

Ag: antigen recognized by scFv antibody